

(I)  $Z_1 \sim Z_2 - X_1 - X_2 - X_3 - X_4 - X_5 - X_6 - X_7 - X_8 - X_9 - X_{10} - Z_3 \sim Z_4$

or a pharmaceutically-acceptable salt thereof, wherein:

$Z_1$  is R-C(O)-NR- or RRN-;

$Z_2$  is an optional 1 to 5 residue peptide or peptide analog;

$X_1$  is any amino acid residue;

$X_2$  is any amino acid residue;

$X_3$  is a hydrophobic residue or a hydroxyl-substituted aliphatic residue;

$X_4$  is any amino acid residue;

$X_5$  is a hydrophobic residue or Gly;

$X_6$  is a hydrophobic or a hydrophilic residue;

$X_7$  is Gly, an amide-substituted polar residue or a hydrophobic residue;

$X_8$  is any amino acid residue;

$X_9$  is an aliphatic residue;

$X_{10}$  is any amino acid residue;

$Z_3$  is an optional 1 to 5 residue peptide or peptide analog;

$Z_4$  is -C(O)OR or -C(O)NRR;

each R is independently hydrogen, (C<sub>1</sub>-C<sub>6</sub>) alkyl, (C<sub>2</sub>-C<sub>6</sub>) alkenyl, (C<sub>2</sub>-C<sub>6</sub>) alkynyl or (C<sub>6</sub>-C<sub>14</sub>) aryl;

each "-" between residues  $X_1$  through  $X_{10}$ ,  $Z_2$  and  $X_1$  and  $X_{10}$  and  $Z_3$

independently represents an amide linkage, a substituted amide linkage or an isostere of an amide linkage; and

each "~" represents a bond.

Substitute the following for claim 4:

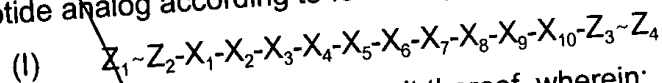
4. (once amended) The compound of claim 1 wherein the mimic comprises a mimic of a chaperone G<sub>1</sub> beta-strand with at least two alternating hydrophobic amino acid residues which exhibits antibacterial activity against a Gram-negative bacterium.

Substitute the following for claim 8:

8. (once amended) The compound of claim 1 wherein the mimic comprises a mimic of an amino terminal motif of a pilus subunit selected from the group consisting of SEQ ID NO: 2, SEQ ID NO: 3, SEQ ID NO: 4, SEQ ID NO: 5, SEQ ID NO: 6, SEQ ID NO: 7, SEQ ID NO: 8, SEQ ID NO: 9, SEQ ID NO: 10, SEQ ID NO: 11, SEQ ID NO: 13, SEQ ID NO: 14, SEQ ID NO: 15, SEQ ID NO: 16, SEQ ID NO: 17, SEQ ID NO: 18, SEQ ID NO: 19, SEQ ID NO: 20, SEQ ID NO: 21, SEQ ID NO: 22, SEQ ID NO: 23, SEQ ID NO: 24, SEQ ID NO: 25, SEQ ID NO: 26, SEQ ID NO: 27, SEQ ID NO: 28 and SEQ ID NO: 29.

Substitute the following from claim 12:

12. (once amended) The compound of claim 1 which is a 10-20 residue peptide or peptide analog according to formula (I):



or a pharmaceutically-acceptable salt thereof, wherein:

$Z_1$  is R-C(O)-NR- or RRN-;

$Z_2$  is an optional 1 to 5 residue peptide or peptide analog;

$X_1$  is any amino acid residue;

$X_2$  is any amino acid residue;

$X_3$  is a hydrophobic residue or a hydroxyl-substituted aliphatic residue;

$X_4$  is any amino acid residue;

$X_5$  is a hydrophobic residue or Gly;

$X_6$  is a hydrophobic or a hydrophilic residue;

$X_7$  is Gly, an amide-substituted polar residue or a hydrophobic residue;

$X_8$  is any amino acid residue;

$X_9$  is an aliphatic residue;

$X_{10}$  is any amino acid residue;

$Z_3$  is an optional 1 to 5 residue peptide or peptide analog;

$Z_4$  is -C(O)OR or -C(O)NRR;

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each R is independently hydrogen,  $\text{C}_1\text{-C}_6$  alkyl,  $\text{C}_2\text{-C}_6$  alkenyl,  $\text{C}_2\text{-C}_6$  alkynyl or  $\text{C}_6\text{-C}_{14}$  aryl;

each "-" between residues  $X_1$  through  $X_{10}$ ,  $Z_2$  and  $X_1$  and  $X_{10}$  and  $Z_3$  independently represents an amide linkage, a substituted amide linkage or an isostere of an amide linkage; and

each "~" represents a bond.

Substitute the following for claim 14:

14. (once amended) The compound of claim 13 which is selected from the group consisting of SEQ ID NO: 2, SEQ ID NO: 3, SEQ ID NO: 4, SEQ ID NO: 5, SEQ ID NO: 6, SEQ ID NO: 7, SEQ ID NO: 8, SEQ ID NO: 9, SEQ ID NO: 10, SEQ ID NO: 11, SEQ ID NO: 13, SEQ ID NO: 14, SEQ ID NO: 15, SEQ ID NO: 16, SEQ ID NO: 17, SEQ ID NO: 18, SEQ ID NO: 19, SEQ ID NO: 20, SEQ ID NO: 21, SEQ ID NO: 22, SEQ ID NO: 23, SEQ ID NO: 24, SEQ ID NO: 25, SEQ ID NO: 26, SEQ ID NO: 27, SEQ ID NO: 28 and SEQ ID NO: 29.

Please cancel claims ~~3, 18~~ and 22-135.

In the Drawings:

Please replace the drawings originally submitted with the formal drawings enclosed.

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